

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/SSO/662
Source: IFWP
Date Processed by STIC: 6/14/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 06/14/2006

PATENT APPLICATION: US/10/550,662

TIME: 10:00:58

Input Set : A:\8030-364US1.TXT

Output Set: N:\CRF4\06142006\J550662.raw

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4 <110> APPLICANT: Sylvia Helen Duncan
5      Harry James Flint
6      Rowett Research Institute
8 <120> TITLE OF INVENTION: Lactic Acid Utilising Bacteria And Their
9      Therapeutic Use
11 <130> FILE REFERENCE: 08830-0364US1
13 <140> CURRENT APPLICATION NUMBER: 10/550,662
C--> 14 <141> CURRENT FILING DATE: 2005-09-26
16 <150> PRIOR APPLICATION NUMBER: GB0307026.5
17 <151> PRIOR FILING DATE: 2003-09-27
19 <160> NUMBER OF SEQ ID NOS: 10
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 20
25 <212> TYPE: DNA
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: Primer
31 <400> SEQUENCE: 1
32 agagtttgat cmtggctcag                                20
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 21
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Primer
42 <400> SEQUENCE: 2
43 acggctacct tgttacgact t                                21
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 18
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Primer
53 <400> SEQUENCE: 3
54 cagcmgccgc ggtaatwc                                    18
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 18
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: Primer
64 <400> SEQUENCE: 4

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65 gwattaccgc ggckgctg 18
67 <210> SEQ ID NO: 5
68 <211> LENGTH: 20
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Primer
75 <400> SEQUENCE: 5
76 aaactcaaaak gaattgacgg 20
78 <210> SEQ ID NO: 6
79 <211> LENGTH: 20
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Primer
86 <400> SEQUENCE: 6
87 ccgtcaattc mtttragttt 20
89 <210> SEQ ID NO: 7
90 <211> LENGTH: 1450
91 <212> TYPE: DNA
92 <213> ORGANISM: Eubacterium hallii (Strain D6 1L/1)
94 <400> SEQUENCE: 7
95 gatgaacgct ggcggcgtgc ctaacactgc aagtcgaacg aagcacctta cctgattctt 60
96 cggatgaagg tctggtgact gagtggcgga cgggtgagta acgcgtgggt aacctgccct 120
97 gtacagggggg ataacagttg gaaacggctg ctaataccgc ataagcgac gagaggacat 180
98 cctcttgtgt gaaaaactcc ggtggtacag gatggggccg cgtctgatta gctggttggc 240
99 agggtaacgg cctaccaagg cgacgatcag tagccggtct gagaggatga acggccacat 300
100 tggaaactgag acacggtcca actcatacgg gaggcagcag tggggaatat tgcacaatgg 360
101 gggaaaccct gatgcagcaa cgccgcgtga gtgaagaagt atttcggtat gtaaagctct 420
102 atcagcaggg aagataatga cggtagctga ctaagaagct ccggctaaat acgtgccagc 480
103 agccgcggta atacgtatgg agcaagcgtt atccggattt actgggtgta aagggtgcgt 540
104 aggtggcagt gcaagtcaga tgtgaaaggc cggggctcaa ccccgagct gcatttgaaa 600
105 ctgcatagct agagtacagg agaggcaggc ggaattccta gtgtagcggg gaaatgcgta 660
106 gatattagga ggaacaccag tggcgaaggc ggcctgctgg actgttactg aactgaggc 720
107 acgaaagcgt ggggagcaaa caggattaga taccctggtg gtccacgccg taaacgatga 780
108 atactaggtg tcggggccgt ataggcttcg gtgccgtcgc aaacgcagta agtattccac 840
109 ctggggagta cgttcgcaag aatgaaactc aaaggaattg acggggaccc gcacaagcgg 900
110 tggagcatgt ggtttaattc gaagcaacgc gaagaacctt accaggtctt gacatccttc 960
111 tgaccactcc gtaatgggag tcttccttcg ggacagaaga gacaggtggg gcatggttgt 1020
112 cgtcagctcg tgtcgtgaga tgttgggtta agtcccgcga cgagcgcaac ccctatcttc 1080
113 agtagccagc aggtaaaggct gggcactctg gagagactgc cagggataac ctggaggaag 1140
114 gtggggacga cgtcaaatca tcatgccctt tatgatctgg gcgacacacg tgctacaatg 1200
115 gcggtcacaa agtgaggcga acctgcgagg gggagcaaac cacaaaaagg ccgtcccagt 1260
116 tcggactgta gtctgcaacc cgactacacg aagctggaat cgctagtaat cgcgaaatcag 1320
117 aatgtcgcgg tgaatacgtt cccgggtctt gtacacaccg cccgtcacac catgggagtc 1380
118 ggaaatgccc gaagccagtg acccaaccat atggaggagg ctgtcgaagg tggagccggg 1440
119 aactgggggtg 1450
121 <210> SEQ ID NO: 8
122 <211> LENGTH: 1458

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123 <212> TYPE: DNA
124 <213> ORGANISM: Eubacterium hallii (Strain SM 6/1)
126 <220> FEATURE:
127 <221> NAME/KEY: misc_feature
128 <222> LOCATION: (1263)...(1263)
129 <223> OTHER INFORMATION: n = a, t, c or g
131 <221> NAME/KEY: misc_feature
132 <222> LOCATION: (1427)...(1427)
133 <223> OTHER INFORMATION: n = a, t, c or g
W--> 135 <221> misc_feature
136 <222> LOCATION: (1455)...(1456)
137 <223> OTHER INFORMATION: n = a, t, c or g
W--> 139 <400> 8
140 gatgaacgct ggcggcggtgc ctaacacatg caagtcgaac gaagcacctt acgagattct 60
141 tcggatgatc gtttggtgac tgagtggcgg acgggtgagt aacgcgtggg taacctgccc 120
142 tgtacagggg gataacagct ggaaacggct gctaataccg cataagcgca cgaggagaca 180
143 tctcctagtg tgaaaaactc cgggtgtaca ggatgggccc gcgtctgatt agctggttgg 240
144 cagggtaacg gcctaccaag gcaacgatca gtagccggtc tgagaggatg aacggccaca 300
145 ttgggaactg gacacggctc aagtcctac gggaggcagc agtggggaat attgcacaat 360
146 ygggyaaaac ctgatgcagc aacgcccgct gagtgaagaa gtatttcggt atgtaaagct 420
147 ctatcagcag ggaagataat gacgggtacct gactaagaag ctccggctaa atacgtgcc 480
148 gcagcccgcg taatagatat ggagcaagcg ttatccggat ttactgggtg taaaggggtg 540
149 gtaggtggca gtgcaagtca gatgtgaaag gccgggggctc aaccccggag ctgcatttga 600
150 aactgcwyrp ctagagtaca ggagaggcag gcggaattcc tagtgtagcg gtgaaatgcg 660
151 tagatattag gaggaacacc agtggcgaa ggcgcctgct ggactgttac tgacactgag 720
152 gcacgaaagc gtggggagca aacaggatta gataccctgg tagtccacgc cgtaaaccat 780
153 gaatactagg tgcgggggccc gtataggctc cgggtgccgc gctaaccgag taagtattcc 840
154 acctggggag tacgttcgca agaatagaac tcaaaggaat tgacggggac ccgcacaagc 900
155 ggtggagcat gtggtttaat tcgaagcaac gcgaagaacc ttaccaggtc ttgacatcct 960
156 tctgaccgca ccttaatcgg tgctttcctt cgggacagaa gagacagggtg gtgcatgggt 1020
157 gtcgtcagct cgtgtcgtga gatgttgggt taagtccgc aacgagcgca acccctatct 1080
158 tcagtagcca gcaggtaagg ctgggcactc tggagagact gccagggata acctggagga 1140
159 aggtggggac gacgtcaaat catcatgcc cttatgatct gggcgacaca cgtgctacaa 1200
160 tggcggtcac agagtgaggc gaaccgcga gggggagcaa accacaaaaa ggccgtccca 1260
W--> 161 gtncggactg tagtctgcaa cccgactaca cagaagctgg aatcgctagt aatcgcgaa 1320
162 cagaatgtcg cggatgaatac gttcccgggt cttgtacaca ccgcccgtca caccatggga 1380
W--> 163 gtcggaaatg cccgaagcca gtgacccaac ctttatgaag gaagccngtc caaggttgaa 1440
W--> 164 cccgttaact ggggnntt 1458
166 <210> SEQ ID NO: 9
167 <211> LENGTH: 1477
168 <212> TYPE: DNA
169 <213> ORGANISM: Ruminococcus gnavus
171 <400> SEQUENCE: 9
172 gagtttgatc ctggctcagg atgaacgctg gcggcggtgcc taacacatgc aagtcgaacg 60
173 aggtatattg aattgaagtt ttcggatgga tttcaatgat accgagtggc ggacgggtga 120
174 gtaacgcgtg ggtaacctgc ctcatacagg gggataacgg ttagaaatga ctgctaatac 180
175 cgcataagcg cacagtaccg catggtacgg tgtgaaaaac tccggtggta tgagatggac 240
176 ccgcgtctga ttagctagtt ggtggggtaa cggcccacca aggcgacgat cagtagccga 300
177 cctgagaggg tgaccggcca cattgggact gagacacggc ccagactcct acgggaggga 360

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178 gcagtggggg atattgcaca atggaggaaa ctctgatgca ggcagccgc gtgagtgaag 420
179 aagtatttcg gtatgtaaag ctctatcagc agggaagaaa atgacggtac ctgactaaga 480
180 agccccggct aactacgtgc cagcagccgc ggtaatacgt agggggcaag cgttatccgg 540
181 atttactggg tgtaaagggg gcgtagacgg cgacgcaagt ctgaagtga ataccggggc 600
182 tcaacctggg aactgctttg gaaactgtgt tgctagagtg ctggagaggt aagcgggaatt 660
183 cctagtgtag cggtgaaatg cgtagatatt aggaagaaca ccagtggcga aggcggctta 720
184 ctggacagta actgacgttg aggtcgaag gcgtggggag caaacaggat tagataacct 780
185 ggtagtccac gccgtaaacg atgaatacta ggtgttggtg agcaaagctc atcggtgccg 840
186 ccgcaaacgc aataagtatt ccacctgggg agtacgttcg caagaatgaa actcaaagga 900
187 attgacgggg acccgacaaa gcggtggagc atgtggttta attcgaagca acgcgaagaa 960
188 ccttaccaaa tcttgacatc cctctgaaaa ryccyttaat cggrrtcctc cttcgggaca 1020
189 gaggtgacag gtggtgcatg gttgtcgta gctcgtgtcg tgagatgttg gggttaagtcc 1080
190 cgcaacgagc gcaacccta ttgtcagtag ccagcaggtg aagctgggca ctctgatgag 1140
191 actgccaggg ataacctgga ggaagggtgg gatgacgtca aatcatcatg ccccttatga 1200
192 tttgggctac acacgtgcta caatggcgta aacaaagaga agcagagctg cgagggggag 1260
193 caaatctcaa aaataacgtc tcagttcgga ttgtagtctg caactcgact acatgaagct 1320
194 ggaatcgcta gtaatcgag atcagaatgc tgcggtgaat acgttcccgg gtcttgatac 1380
195 caccgcccgt cacaccatgg gagtcggaaa tgcccgaagc cagtgaaccc aatgcgaaag 1440
196 cagggaagctg tgaagggcag gtctgataac tggggtg 1477
198 <210> SEQ ID NO: 10
199 <211> LENGTH: 1493
200 <212> TYPE: DNA
201 <213> ORGANISM: Clostridium indolis
203 <400> SEQUENCE: 10
204 agagtttgat cctggctcag gatgaacgct ggcggcgtgc ttaacacatg caagtcgaac 60
205 gaaacacctt atttgatttt cttcggaact gaagatttgg tgattgagtg gcggacgggt 120
206 gagtaacgcg tgggtaacct gccctgtaca gggggataac agtcagaaat gactgctaata 180
207 accgcataag accacagcac cgcattggtc aggggtaaaa actccggtgg tacaggatgg 240
208 acccgctgtc gattagctgg ttggtgaggt aacggctcac caaggcgacg atcagtagcc 300
209 ggcttgagag agtgaacggc cacattggga ctgagacacg gcccacactc ctacgggagg 360
210 cagcagtggg gaatatgca caatggggga aacctgatg cagcagacgc gcgtgagtga 420
211 agaagtatct cggtagtaa agctctatca gcagggaaga aaatgacggt acctgactaa 480
212 gaagccccgg ctaactacgt gccagcagcc gcgtaatac gtagggggca agcgttatcc 540
213 ggaattactg ggtgtaaagg gtgcgtaggt ggtatggcaa gtcagaagtg aaaacccagg 600
214 gcttaactct gggactgctt ttgaaactgt cagactggag tgcaggagag gtaagcggaa 660
215 ttcttagtgt agcggtgaaa tgcgtagata ttaggaggaa catcagtggc gaaggcggct 720
216 tactggactg aaactgacac tgaggcacga aagcgtgggg agcaaacagg attagatacc 780
217 ctggtagtcc acgcgtaaa cgatgaatac taggtgtcgg gcccgtagag gcttcggtgc 840
218 cgagccaac gcagtaagta ttccacctgg ggagtacgtt cgcaagaatg aactcaaagg 900
219 aattgacggg gacccgcaca agcgggtggag catgtggttt aattcgaagc aacgcgaaga 960
220 accttacctg gtcttgacat ccttctgacc ggtccttaac cggacctttc cttcgggaca 1020
221 ggagtgcag gtggtgcatg gttgtcgta gctcgtgtcg tgagatgttg gggttaagtcc 1080
222 cgcaacgagc gcaacccta tctttagtag ccagcatata agtggggac tctagagaga 1140
223 ctgccaggga taactggag gaaggtgggg acgacgtcaa atcatcatgc ccttatgac 1200
224 cagggtctaca cacgtgctac aatggcgtaa acaggggaa gcagcctcgt gagagtgagc 1260
225 aaatccaaa aataacgtct cagttcggat tgtagtctgc aactcgacta catgaagctg 1320
226 gaatcgctag taatcgcgaa tcagaatgtc gcggtgaata cgttcccggg tcttgtaac 1380
227 accgcccgtc acaccatggg agtcagtaac gcccgaaagc agtgacccaa ccgtaaggag 1440
228 gagctgccga agcgggaccg ataactgggg tgaagtcgta accaggtagc cgt 1493

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/550,662

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TIME: 10:00:59

Input Set : A:\8830-364US1.TXT
Output Set: N:\CRF4\06142006\J550662.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:8; N Pos. 1263,1427,1455,1456

VERIFICATION SUMMARY

DATE: 06/14/2006

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Input Set : A:\8830-364US1.TXT

Output Set: N:\CRF4\06142006\J550662.raw

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:135 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:139 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:161 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:1260
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:1380
L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:1440